

## SEQUENCE LISTING

- <110> Dalton, John Andrews, Stuart
- <120> Vaccine Containing a Thiol Protease
- <130> 1181-281
- <140> 10/ 620,451
- <141> 2003-07-17
- <150> US 08/ 424,361
- <151> 1995-05-25
- <150> PCT/ GB93/ 02172
- <151> 1993-10-21
- <160> 23
- <170> PatentIn version 3.2
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- Val Lys Asp Gln 20
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- Lys Asp Gln
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Gln Gly Asn Cys Xaa Xaa Cys Trp Ala Phe Ser Thr Thr Gly Thr Met
1
                                    10
                                                         15
gag gga caa tat atg aaa aac gaa aaa act agt att tca ttc tct qag
                                                                       96
Glu Gly Gln Tyr Met Lys Asn Glu Lys Thr Ser Ile Ser Phe Ser Glu
            20
                                                     30
caa caa ctg gtc gat tgt agc ggt cct tgg gga aat aat ggt tgc agt
                                                                      144
Gln Gln Leu Val Asp Cys Ser Gly Pro Trp Gly Asn Asn Gly Cys Ser
        35
                            40
                                                 45
ggt gga ttg atg gaa aat gct tac caa tat ttg aaa caa ttt gga ttg
                                                                      192
Gly Gly Leu Met Glu Asn Ala Tyr Gln Tyr Leu Lys Gln Phe Gly Leu
    50
                        55
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gaa acc gaa tcc tct tat ccg tac acg gct gtg gaa ggt cag tgt cga
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Glu Thr Glu Ser Ser Tyr Pro Tyr Thr Ala Val Glu Gly Gln Cys Arg
65
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tac aat agg cag ttg gga gtt gcc aaa gtg acc ggc tac tat act gtg
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Tyr Asn Arg Gln Leu Gly Val Ala Lys Val Thr Gly Tyr Tyr Thr Val
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                                     90
                                                          95
cat tct ggc agt gag gta gaa ttg aaa aat cta gtc ggt tcc gaa gga
                                                                       336
His Ser Gly Ser Glu Val Glu Leu Lys Asn Leu Val Gly Ser Glu Gly
            100
                                 105
                                                      110
cct gcc gcg atc gct gtg gat gtg gaa tct gac ttc atg atg tac agg
                                                                       384
Pro Ala Ala Ile Ala Val Asp Val Glu Ser Asp Phe Met Met Tyr Arg
        115
                             120
                                                 125
agt ggt att tat cag agc caa act tgt tta ccg ttc gct ctg aat cat
                                                                       432
Ser Gly Ile Tyr Gln Ser Gln Thr Cys Leu Pro Phe Ala Leu Asn His
    130
                         135
                                             140
gca gtc ttg tct gtc ggt tat gga aca cag gat ggt act gnt t
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                                                         15
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            20
                                 25
                                                     30
Gln Gln Leu Val Asp Cys Ser Gly Pro Trp Gly Asn Asn Gly Cys Ser
        35
                             40
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Gly Gly Leu Met Glu Asn Ala Tyr Gln Tyr Leu Lys Gln Phe Gly Leu 50 55 Glu Thr Glu Ser Ser Tyr Pro Tyr Thr Ala Val Glu Gly Gln Cys Arg 65 70 75 80 Tyr Asn Arg Gln Leu Gly Val Ala Lys Val Thr Gly Tyr Tyr Thr Val 85 90 95 His Ser Gly Ser Glu Val Glu Leu Lys Asn Leu Val Gly Ser Glu Gly 100 105 110 Pro Ala Ala Ile Ala Val Asp Val Glu Ser Asp Phe Met Met Tyr Arg 115 120 125 Ser Gly Ile Tyr Gln Ser Gln Thr Cys Leu Pro Phe Ala Leu Asn His 130 135 140 Ala Val Leu Ser Val Gly Tyr Gly Thr Gln Asp Gly Thr Xaa 145 150 155 <210> 19 <211> 477 <212> DNA <213> Fasciola hepatica <220> <221> misc feature <222> (1)..(477)<223> n = any nucleotide <220> <221> CDS <222> (1)..(477) <400> 19 cat caa gaa gcc cnn ggc tct tgt tgg gnt ttc tca aca aca ggt gct 48 His Gln Glu Ala Xaa Gly Ser Cys Trp Xaa Phe Ser Thr Thr Gly Ala 1 10 15 atg gaa gga cag tat atg aaa aac caa aga act agt att tca tnc tct 96 Met Glu Gly Gln Tyr Met Lys Asn Gln Arg Thr Ser Ile Ser Xaa Ser 20 25 30 gag caa caa ctg gtc gat tgt agc cgt gat ttt ggc aat tat ggt tgt 144 Glu Gln Gln Leu Val Asp Cys Ser Arg Asp Phe Gly Asn Tyr Gly Cys

35 40 45

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aat ggt gga cta atg gaa aat gca tac gaa tat ttg aaa cga ttt gga
                                                                       192
Asn Gly Gly Leu Met Glu Asn Ala Tyr Glu Tyr Leu Lys Arg Phe Gly
    50
                         55
ttg gaa acc gag tct tct tat cct tac agg gct gtg gaa gga caa tgt
                                                                       240
Leu Glu Thr Glu Ser Ser Tyr Pro Tyr Arg Ala Val Glu Gly Gln Cys
65
                     70
                                         75
                                                              80
cga tac aac gag cag ttg gga gtt gcc aaa gtg act agc tac tat acg
                                                                       288
Arg Tyr Asn Glu Gln Leu Gly Val Ala Lys Val Thr Ser Tyr Tyr Thr
                85
                                     90
                                                          95
gta cat tct gga gat gag gta gaa ttg caa aat cta gtc ggt gcc gaa
                                                                       336
Val His Ser Gly Asp Glu Val Glu Leu Gln Asn Leu Val Gly Ala Glu
            100
                                 105
                                                     110
gga cct gct gcg gtc gct ttg gat gtg gag tca gac ttc atg atg tac
                                                                       384
Gly Pro Ala Ala Val Ala Leu Asp Val Glu Ser Asp Phe Met Met Tyr
        115
                             120
                                                 125
agg agt ggt att tat cag agc caa act tgt tca ccg gat cgt ttg aac
                                                                       432
Arg Ser Gly Ile Tyr Gln Ser Gln Thr Cys Ser Pro Asp Arg Leu Asn
    130
                        135
                                             140
cat gga gtg ttg nct gtc gnt tat gga acn cag ggt ggt nct cnc
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- <223> The 'Xaa' at location 149 stands for Thr, Ala, Pro, or Ser.

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Asn Gly Gly Leu Met Glu Asn Ala Tyr Glu Tyr Leu Lys Arg Phe Gly 50 60

Leu Glu Thr Glu Ser Ser Tyr Pro Tyr Arg Ala Val Glu Gly Gln Cys 70 75 80

Arg Tyr Asn Glu Gln Leu Gly Val Ala Lys Val Thr Ser Tyr Tyr Thr 85 90 95

Val His Ser Gly Asp Glu Val Glu Leu Gln Asn Leu Val Gly Ala Glu
100 105 110

Gly Pro Ala Ala Val Ala Leu Asp Val Glu Ser Asp Phe Met Met Tyr 115 120 125

Arg Ser Gly Ile Tyr Gln Ser Gln Thr Cys Ser Pro Asp Arg Leu Asn 130 135

His Gly Val Leu Xaa Val Xaa Tyr Gly Thr Gln Gly Gly Xaa Xaa

145 150 155

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                                                          15
gga caa tat atg aaa aac gaa aaa act agt ntt tca ncc tct gag caa
                                                                        96
Gly Gln Tyr Met Lys Asn Glu Lys Thr Ser Xaa Ser Xaa Ser Glu Gln
            20
                                 25
                                                      30
caa ctg gtc gat tgt agc ggt cct tgg gga aat aat ggt tgc agt ggt
                                                                       144
Gln Leu Val Asp Cys Ser Gly Pro Trp Gly Asn Asn Gly Cys Ser Gly
        35
                             40
                                                 45
gga ttg atg gaa aat gct tac caa tat tta aaa caa ttt gga ttg gaa
                                                                       192
Gly Leu Met Glu Asn Ala Tyr Gln Tyr Leu Lys Gln Phe Gly Leu Glu
    50
                         55
                                             60
acc gaa tcc tct tat ccg tac acg gct gtg gaa ggt cag tgt cga tac
Thr Glu Ser Ser Tyr Pro Tyr Thr Ala Val Glu Gly Gln Cys Arg Tyr
65
                    70
                                                              80
                                         75
aat agg cag ttg gga gtt gcc aaa gtg act ggc tac tat act gtg cat
                                                                       288
Asn Arg Gln Leu Gly Val Ala Lys Val Thr Gly Tyr Tyr Thr Val His
                85
                                     90
                                                          95
tct ggc agt gag gca gga ttg aaa aat cta gtc ggt tcc gaa gga cct
                                                                       336
Ser Gly Ser Glu Ala Gly Leu Lys Asn Leu Val Gly Ser Glu Gly Pro
            100
                                 105
                                                     110
gcc gcg atc gct gtg gat gtg gaa tct gac ttc atg atg tac agg agt
                                                                       384
Ala Ala Ile Ala Val Asp Val Glu Ser Asp Phe Met Met Tyr Arg Ser
        115
                             120
                                                 125
ggt att tat cag anc caa act tgt tta ccg ttc gct ttg aat cat gca
                                                                       432
Gly Ile Tyr Gln Xaa Gln Thr Cys Leu Pro Phe Ala Leu Asn His Ala
    130
                        135
                                             140
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Val Leu Xaa Val Asp Tyr Gly Thr Gln Asp Gly Xaa Xaa
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       PRT
<213>
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                                 25
                                                     30
Gln Leu Val Asp Cys Ser Gly Pro Trp Gly Asn Asn Gly Cys Ser Gly
        35
                             40
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472

Gly Leu Met Glu Asn Ala Tyr Gln Tyr Leu Lys Gln Phe Gly Leu Glu

50 55 60

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Asn Arg Gln Leu Gly Val Ala Lys Val Thr Gly Tyr Tyr Thr Val His
85 90 95

Ser Gly Ser Glu Ala Gly Leu Lys Asn Leu Val Gly Ser Glu Gly Pro 100 105 110

Ala Ala Ile Ala Val Asp Val Glu Ser Asp Phe Met Met Tyr Arg Ser 115 120 125

Gly Ile Tyr Gln Xaa Gln Thr Cys Leu Pro Phe Ala Leu Asn His Ala 130 135 140

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<212> PRT

<213> Fasciola hepatica

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